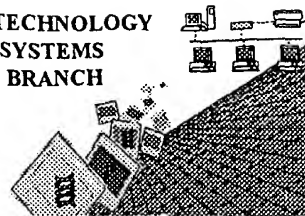


0300
BIOTECHNOLOGY
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RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/896,896
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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



OIPE

RAW SEQUENCE LISTING

DATE: 02/07/2002

PATENT APPLICATION: US/09/896,896

TIME: 13:36:42

Input Set : A:\31-17US Sequence listing.txt

Output Set: N:\CRF3\02072002\I896896.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: MAXYGEN APS
5 <120> TITLE OF INVENTION: N-TERMINALLY EXTENDED POLYPEPTIDES
7 <130> FILE REFERENCE: 0217us210
W--> 9 <140> CURRENT APPLICATION NUMBER: US/09/896,896
W--> 0 <160> NUMBER OF SEQ ID NOS:
9 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

see next page

09/896,896 2

SEQUENCE LISTING

<110> MAXYGEN APS

<120> N-TERMINALLY EXTENDED POLYPEPTIDES

<130> 0217us210

<170> PatentIn Ver. 2.1

<210> 1

→ <1607> this is a mandatory
numeric identifier and
needs a response

FYI: 1/3 sequences in
submitted file

FYI

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/896,896

DATE: 02/07/2002

TIME: 13:36:44

Input Set : A:\31-17US Sequence listing.txt

Output Set: N:\CRF3\02072002\I896896.raw

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L:0 M:282 W: Numeric Field Identifier Missing, <160> is required.
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:156 M:283 W: Missing Blank Line separator, <220> field identifier
L:160 M:283 W: Missing Blank Line separator, <400> field identifier
L:392 M:283 W: Missing Blank Line separator, <220> field identifier
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L:668 M:283 W: Missing Blank Line separator, <400> field identifier
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L:2446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:101
L:0 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (0) Counted (113)